

BS — compared with the EDG1-7 polypeptides (EDG1 (SEQ ID NO:6): accession number M 31210, EDG2 (SEQ ID NO:3): accession number U 80811, EDG3 (SEQ ID NO:7): accession number X 83864, EDG4 (SEQ ID NO:5): accession number AF 011466, EDG5 (SEQ ID NO:8): accession number AF 034780, EDG6 (SEQ ID NO:9): AJ 000479, EDG7 (SEQ ID NO:4): accession number AF 127138). The approximate boundaries of the seven putative transmembrane domains are boxed. Gaps are introduced to optimize the alignment.

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